

				50					55					60
Phe	Phe	Thr	Gly	Pro 65	Met	Ser	Asp	Phe	Lys 70	Asn	Val	Gly	Leu	Val 75
Phe	Val	Asn	Ser	Lys 80	Arg	Asp	Arg	Thr	Lys 85	Ala	Val	Leu	Cys	Met 90
Val	Val	Ala	Gly	Ala 95	Ile	Ala	Ala	Val	Phe 100	His	Thr	Leu	Ile	Ala 105
Tyr	Ser	Asp	Leu	Gly 110	Tyr	Tyr	Ile	Ile	Asn 115	Lys	Leu	His	His	Val 120
Asp	Glu	Ser	Val	Gly 125	Ser	Lys	Thr	Arg	Arg 130	Ala	Phe	Leu	Tyr	Leu 135
Ala	Ala	Phe	Pro	Phe 140	Met	Asp	Ala	Met	Ala 145	Trp	Thr	His	Ala	Gly 150
Ile	Leu	Leu	Lys	His 155	Lys	Tyr	Ser	Phe	Leu 160	Val	Gly	Cys	Ala	Ser 165
Ile	Ser	Asp	Val	Ile 170	Ala	Gln	Val	Val	Phe 175	Val	Ala	Ile	Leu	Leu 180
His	Ser	His	Leu	Glu 185	Cys	Arg	Glu	Pro	Leu 190	Leu	Ile	Pro	Ile	Leu 195
Ser	Leu	Tyr	Met	Gly 200	Ala	Leu	Val	Arg	Cys 205	Thr	Thr	Leu	Cys	Leu 210
Gly	Tyr	Tyr	Lys	Asn 215	Ile	His	Asp	Ile	Ile 220	Pro	Asp	Arg	Ser	Gly 225
Pro	Glu	Leu	Gly	Gly 230	Asp	Ala	Thr	Ile	Arg 235	Lys	Met	Leu	Ser	Phe 240
Trp	Trp	Pro	Leu	Ala 245	Leu	Ile	Leu	Ala	Thr 250	Gln	Arg	Ile	Ser	Arg 255
Pro	Ile	Val	Asn	Leu 260	Phe	Val	Ser	Arg	Asp 265	Leu	Gly	Gly	Ser	Ser 270
Ala	Ala	Thr	Glu	Ala 275	Val	Ala	Ile	Leu	Thr 280	Ala	Thr	Tyr	Pro	Val 285
Gly	His	Met	Pro	Tyr 290	Gly	Trp	Leu	Thr	Glu 295	Ile	Arg	Ala	Val	Tyr 300
Pro	Ala	Phe	Asp	Lys 305	Asn	Asn	Pro	Ser	Asn 310	Lys	Leu	Val	Ser	Thr 315
Ser	Asn	Thr	Val	Thr 320	Ala	Ala	His	Ile	Lys 325	Lys	Phe	Thr	Phe	Val 330
Cys	Met	Ala	Leu	Ser 335	Leu	Thr	Leu	Cys	Phe 340	Val	Met	Phe	Trp	Thr 345

Pro Asn Val Ser	Glu Lys Ile Leu Ile	Asp Ile Ile Gly Val Asp
	350	355 360
Phe Ala Phe Ala	Glu Leu Cys Val Val	Pro Leu Arg Ile Phe Ser
	365	370 375
Phe Phe Pro Val	Pro Val Thr Val Arg	Ala His Leu Thr Gly Trp
	380	385 390
Leu Met Thr Leu	Lys Lys Thr Phe Val	Leu Ala Pro Ser Ser Val
	395	400 405
Leu Arg Ile Ile	Val Leu Ile Ala Ser	Leu Val Val Leu Pro Tyr
	410	415 420
Leu Gly Val His	Gly Ala Thr Leu Gly	Val Gly Ser Leu Leu Ala
	425	430 435
Gly Phe Val Gly	Glu Ser Thr Met Val	Ala Ile Ala Ala Cys Tyr
	440	445 450
Val Tyr Arg Lys	Gln Lys Lys Lys Met	Glu Asn Glu Ser Ala Thr
	455	460 465
Glu Gly Glu Asp	Ser Ala Met Thr Asp	Met Pro Pro Thr Glu Glu
	470	475 480
Val Thr Asp Ile	Val Glu Met Arg Glu	Glu Asn Glu
	485	490

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 <211> 535
 <212> DNA
 <213> Homo sapiens

<220>
 <221> unsure
 <222> 33, 66, 96, 387
 <223> unknown base

<400> 8
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 cggcctattg tcaacctctt tgtttcccg gaccttggtg gcagttctgc 150
 agccacagag gcagtggcga ttttgacagc cacataccct gtgggtcaca 200
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 aagaataacc ccagcaacaa actggtgagc acgagcaaca cagtcacggc 300
 ggccacatc aagaagttca cttcgtctg catggctctg tcaactcagc 350
 tctgtttcgt gatgttttg acaccaacg tgtctgngaa aatcttgata 400
 gacatcatcg gagtggactt tgcctttgca gaactctgtg ttgttcoctt 450

